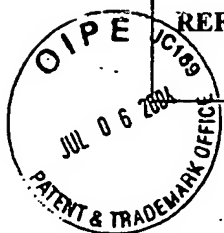


Form PTO-1449 (adapted)	Docket No. 133.00US	Serial No. 10/726,321
REFERENCES CITED BY APPLICANT	First Named Inventor Dear Bing	Customer No. 53,603
	Filing Date 01 December 2003	Group 1753



U.S. PATENT DOCUMENTS

Examiner's Initial		Document Number	Inventor(s)	Class /Subclass	Title	Issue Date or Publ. Date (dd.mm.yy)
an	P1	4,314,343	TOMLINSON	364/498	Method and apparatus for detecting and integrating chromatographic peaks	02 Feb 82
an	P2	4,353,242	HARRI	73/23.1	Multichannel detection and resolution of chromatographic peaks	12 Oct 82
an	P3	4,837,726	HUNKAPILLER	364/498	Quantitation of chromatographic information	06 Jun 89
an	P4	4,865,706	KARGER	204/182.8	High performance microcapillary gel electrophoresis	12 Sep 89
an	P5	5,098,536	ANDERSON	204/180.1	Method of improving signal-to-noise in electropherogram	24 Mar 92
an	P6	5,470,705	GROSSMAN	435/6	Probe composition containing a binding domain and polymer chain and methods of use	28 Nov 95
an	P7	5,502,773	TIBBETTS	382/129	Method and apparatus for automated processing of DNA sequence data	26 Mar 96
an	P8	5,503,994	SHEAR	436/90	System for sample detection with compensation for difference in sensitivity to detection of components moving at different velocities	02 Apr 96
an	P9	5,873,052	SHARAF	702/020	Alignment-based similarity scoring methods for quantifying the differences between related biopolymer sequences	16 Feb 99
an	P10	5,905,192	WIKFORS	073/023.35	Method for identification of chromatographic peaks	18 May 99
an	P11	5,916,747	GILCHRIST	435/006	Method and apparatus for alignment of signals for use in DNA based-calling	29 Jun 99
an	P12	6,502,039	THEILHABER	702/019	Mathematical analysis for the estimation of changes in the level of gene expression	31 Dec 02
an	P13	6,505,125	HO	702/019	Methods and computer software products for multiple probe gene expression analysis	7 Jan 03
an	P14	6,510,391	BALABAN	702/019	Computer software products for nucleic acid hybridization analysis	21 Jan 03
an	P15	6,516,276	GHANDOUR	702/027	Method and apparatus for analysis of data from biomolecular arrays	04 Feb 03

EXAMINER <i>Ch. Koznerola</i>	Date considered <i>01/31/06</i>
*EXAMINER: Initial if reference considered, whether or not citation in conformance with MPEP 609; Draw line through citation if not in conformance and/or not considered. Include copy of this form with next communication to applicant.	

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am	D1	NAVAKATIKYAN, "A Real-Time Algorithm for Quantification of Blood Pressure Waveforms", IEEE Transactions on Biomedical Engineering, Vol. 49, No. 7, July 2002, pgs. 662-670.
am	D2	IZMAILLOV, "Improvement of Base-Calling in Multilane Automated DNA Sequencing by Use of Electrophoretic Calibration Standards, Data Linearization, and Trace Alignment", Electrophoresis, Vol. 22, 2002, pgs. 1906-1914.

EXAMINER	<i>Ch. Meyers</i>	Date considered	<i>01/31/06</i>
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*EXAMINER: Initial if reference considered, whether or not citation in conformance with MPEP 609; Draw line through citation if not in conformance and/or not considered. Include copy of this form with next communication to applicant.